

Unnatural Amino Acid Incorporation in Virus-like Particles

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Supporting Information

1. Expression of Q β (K16M-CCH)₉₀ in M15(pREP4)MA cells.

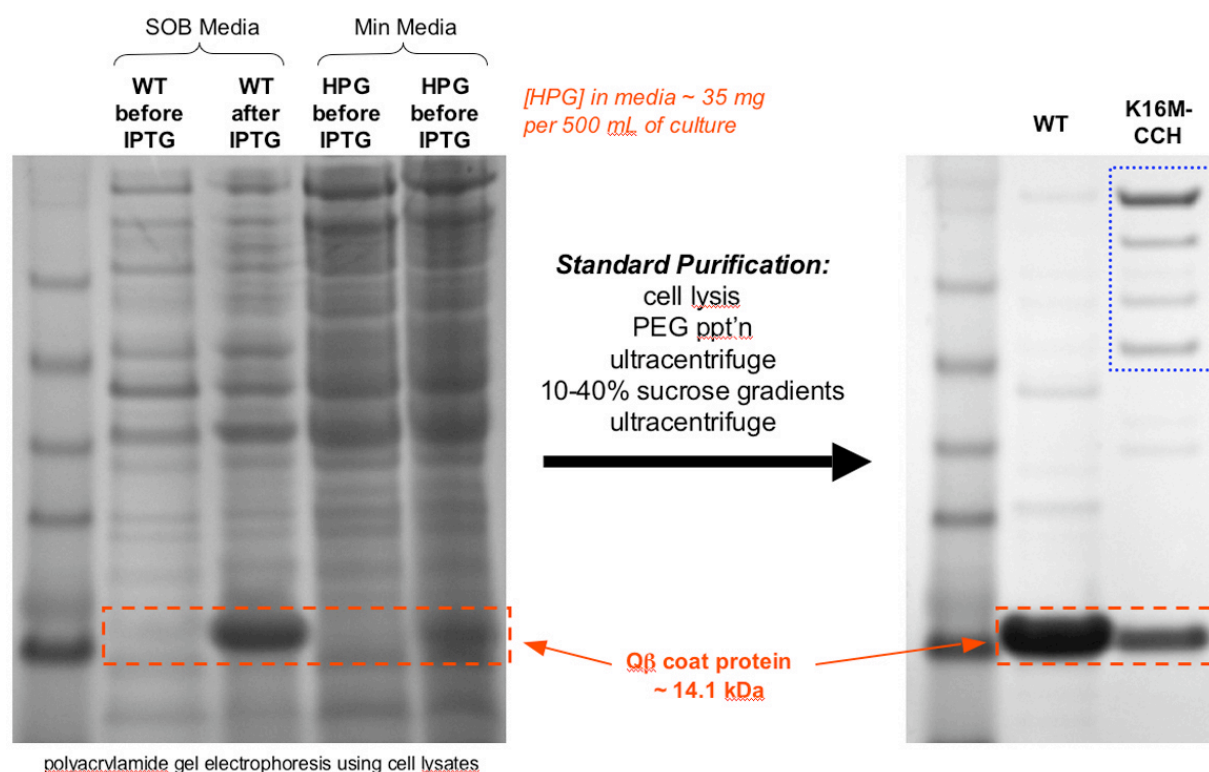


Figure S1. (Left) Gel electrophoresis analyses of cell lysates showing IPTG-induced expression of Q β coat protein under standard (SOB media) and unnatural (Min media) conditions, the latter in the presence of homopropargylglycine **2**. (Right) Gel electrophoresis analysis of purified Q β (K16M-CCH)₉₀ particles. The bands in the blue box are impurities which are removed by a second sucrose gradient ultracentrifugation.

2. Media for expression of unnatural amino acids in M15(pREP4)MA cells.

M9 Minimal Media	25X M9 Salts	19 Amino Acid Mix
500 mL of doubly deionized H ₂ O	320 g Na ₂ HPO ₄ •7H ₂ O	1 L doubly deionized H ₂ O
25 mL 25X M9 salts	75 g KH ₂ PO ₄	2g serine
25 mL 20% (w/v) glucose	12.5 g NaCl	1 g remaining amino acids minus methionine
25 mL 19 Amino Acid Mix	25 g NH ₄ Cl	
2mL 1M MgCl	doubly deionized H ₂ O to 1 L	
0.2 mL 1M CaCl ₂	pH 7.4	
4 mL methionine or 40 mg azidohomoalanine		

3. Characterization of Q β (K16M-CCH)₉₀.

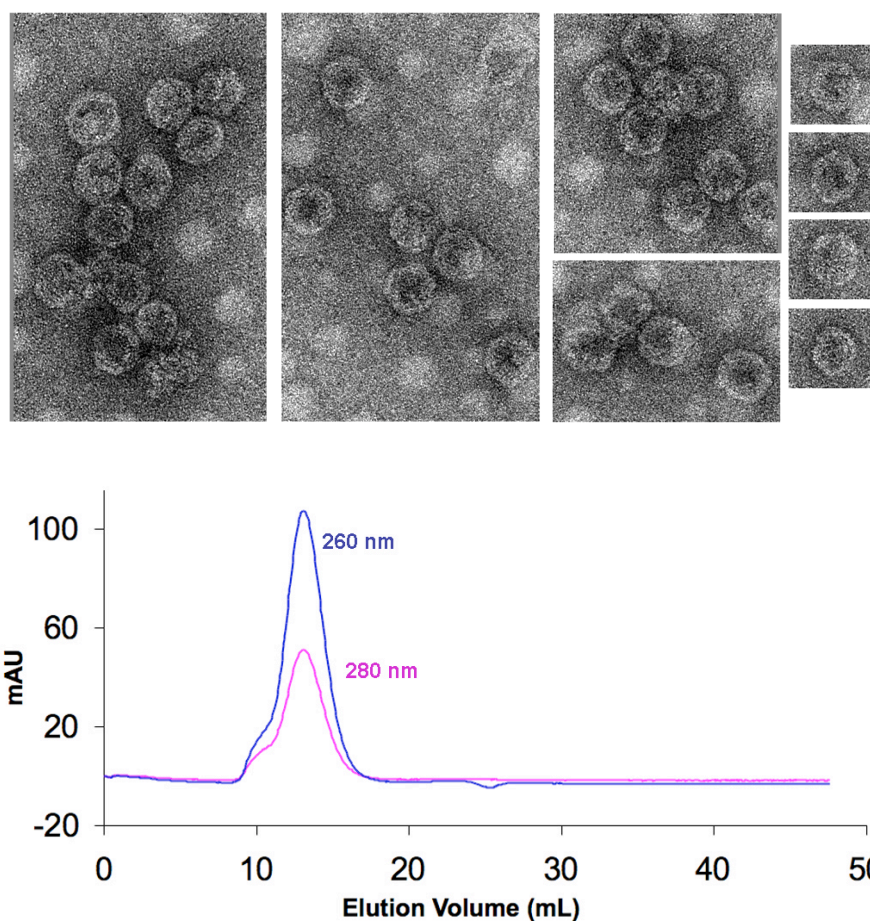


Figure S2. (Top) Negative-stained (1% uranyl acetate) TEM images (45K magnification) of Q β (K16M-CCH)₉₀, showing 28-nm diameter particles. The white circular images are from the background grid and are not due to protein. (Bottom) Size-exclusion chromatography (Superose-6) of Q β (K16M-CCH)₉₀, showing the retention volume and A_{260}/A_{280} ratio characteristic of intact Q β particles.

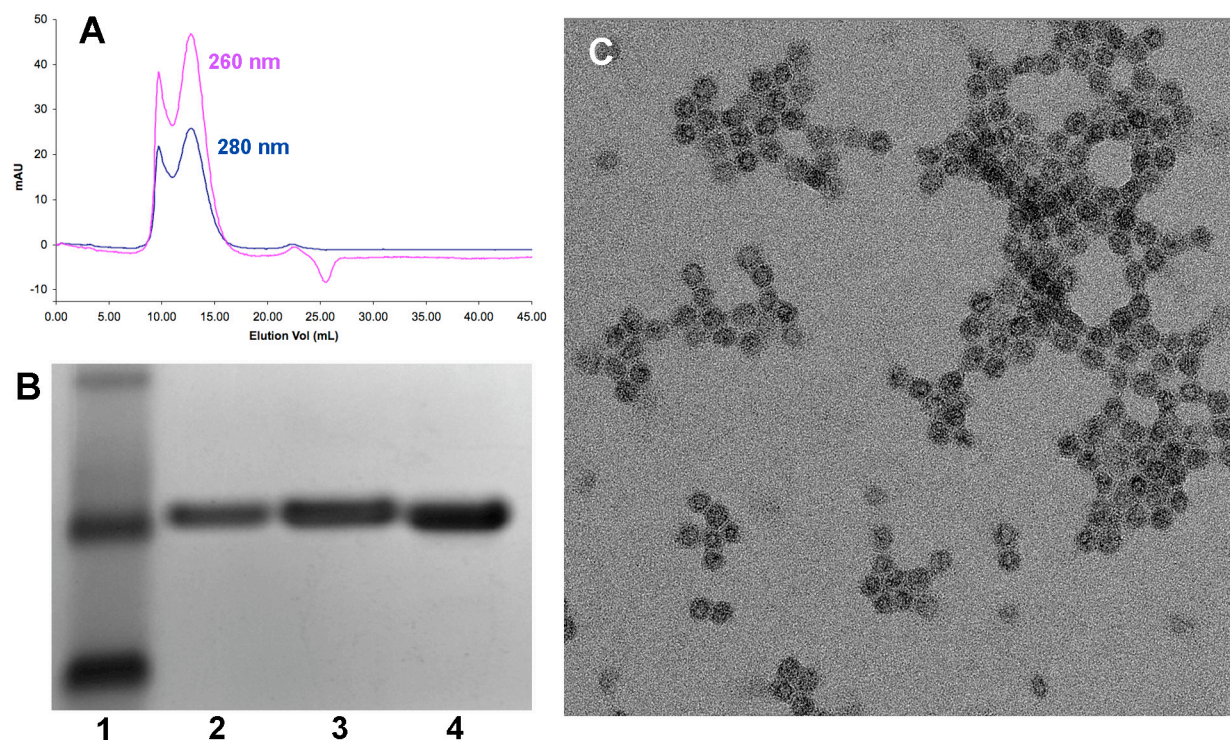
4. Characterization of Q β (K16M-CCH)₉₀ adduct with azide **11**.

Figure S3. (A) Size-exclusion chromatography (Superose-6) of the purified adduct of Q β (K16M-CCH)₉₀ with azide **11**. This material contained a mixture of two fractions eluted from the 10-40% sucrose gradient, comprising a “tight” band of material at the characteristic density of Q β VLPs, and a more diffuse band at slightly higher density. The observation of two fractions in the size-exclusion chromatogram suggests some aggregation of particles under both conditions. (B) Denaturing polyacrylamide gel electrophoresis of the purified adduct in part (A). Lane 1 = ladder; lane 2 = top fraction; lane 3 = bottom fraction; lane 4 = wild-type Q β , unlabeled. (C) Negative-stained (1% uranyl acetate) TEM image (45K magnification) of the purified adduct in part (A), showing intact particles.

Table S1. Reported yields of proteins incorporating unnatural amino acids. Note that many more examples are known, but these comprise most of those that describe isolated yields of significant amounts of protein. DHFR = dihydrofolate reductase; GFP = green fluorescent protein; Pr A = Z-domain of protein A; SD = superoxide dismutase.

Unnatural Amino Acid Incorporated	Protein	Protein yield (mg/L)	Ref.
3,4-dihydroxy-L-phenylalanine	Myoglobin	1	[1]
<i>p</i> -benzoyl-L-phenylalanine	Myoglobin	2	[2]
<i>p</i> -azido-L-phenylalanine	Myoglobin	2	[3]
¹⁵ N labeled <i>p</i> -methoxyphenylalanine	Myoglobin	1	[4]
<i>p</i> -azido-L-phenylalanine	SD	0.8	[5]
azidohomoalanine	DHFR	35	[6]
2-amino-5-hexynoic acid	DHFR	35	[6]
norleucine	DHFR	20	[6]
homoallylglycine	DHFR	10	[7]
<i>p</i> -iodophenylalanine	DHFR	6-18	[8]
<i>p</i> -cyanophenylalanine	DHFR	6-18	[8]
<i>p</i> -ethynylphenylalanine	DHFR	6-18	[8]
<i>p</i> -azidophenylalanine	DHFR	6-18	[8]
2-pyrindinylalanine	DHFR	6-18	[8]
3-pyrindinylalanine	DHFR	6-18	[8]
4-pyrindinylalanine	DHFR	6-18	[8]
<i>p</i> -acetyl-L-phenylalanine	Pr A	3.6	[9]
(2S,3S)-2-aminomethyl pentanoic acid	DHFR	12-22	[10]
trifluoroisoleucine	DHFR	20	[11]
<i>p</i> -nitrophenylalanine	Pr A	2	[12]
homoallylglycine	DHFR	70	[13, 14]
2-amino-5-hexynoic acid	DHFR	8	[13]
L-3-(2-naphthyl)alanine	DHFR	2.2	[15]
<i>p</i> -amino-L-phenylalanine	GFP	10-22	[16]
<i>p</i> -methoxy-L-phenylalanine	GFP	10-22	[16]
<i>p</i> -iodo-L-phenylalanine	GFP	10-22	[16]
<i>p</i> -bromo-L-phenylalanine	GFP	10-22	[16]
L-3-(2-naphthyl)alanine	GFP	10-22	[16]
<i>p</i> -acetyl-phenylalanine	Pr A	3.6-9.2	[17]
<i>o</i> -methyltyrosine	SD	0.6	[18]
α -aminocaprylic acid	SD	0.6	[18]
<i>o</i> -nitrobenzyl cysteine	SD	0.6	[18]
<i>p</i> -iodo-L-phenylalanine	Pr A	3.8-4.6	[19]
<i>p</i> -iodo-L-phenylalanine	T4 Lysozyme	5.7	[19]
N-acetylgalactosamine- α - <i>o</i> -threonine	Myoglobin	2.4	[20]

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